

Figure 1

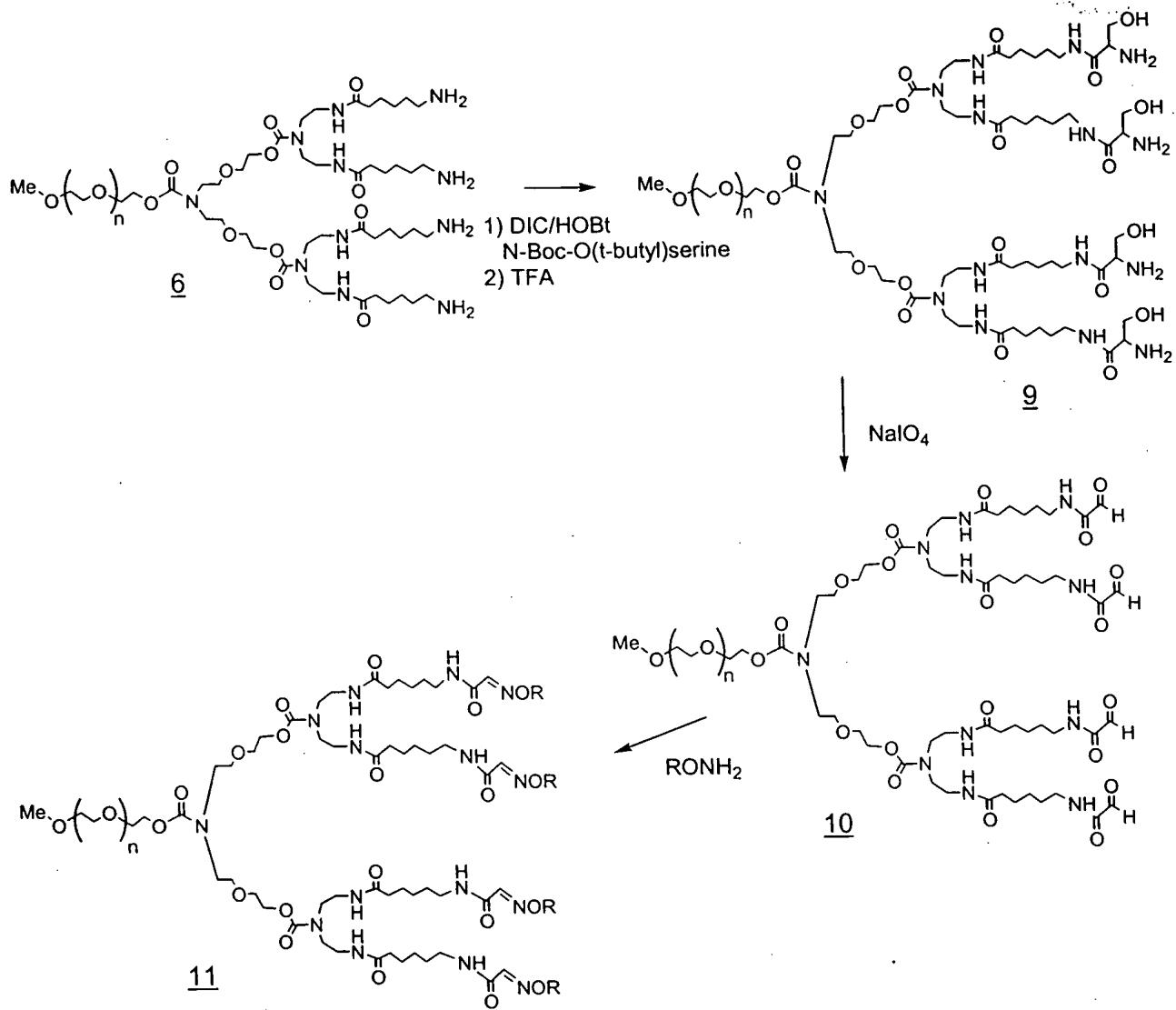
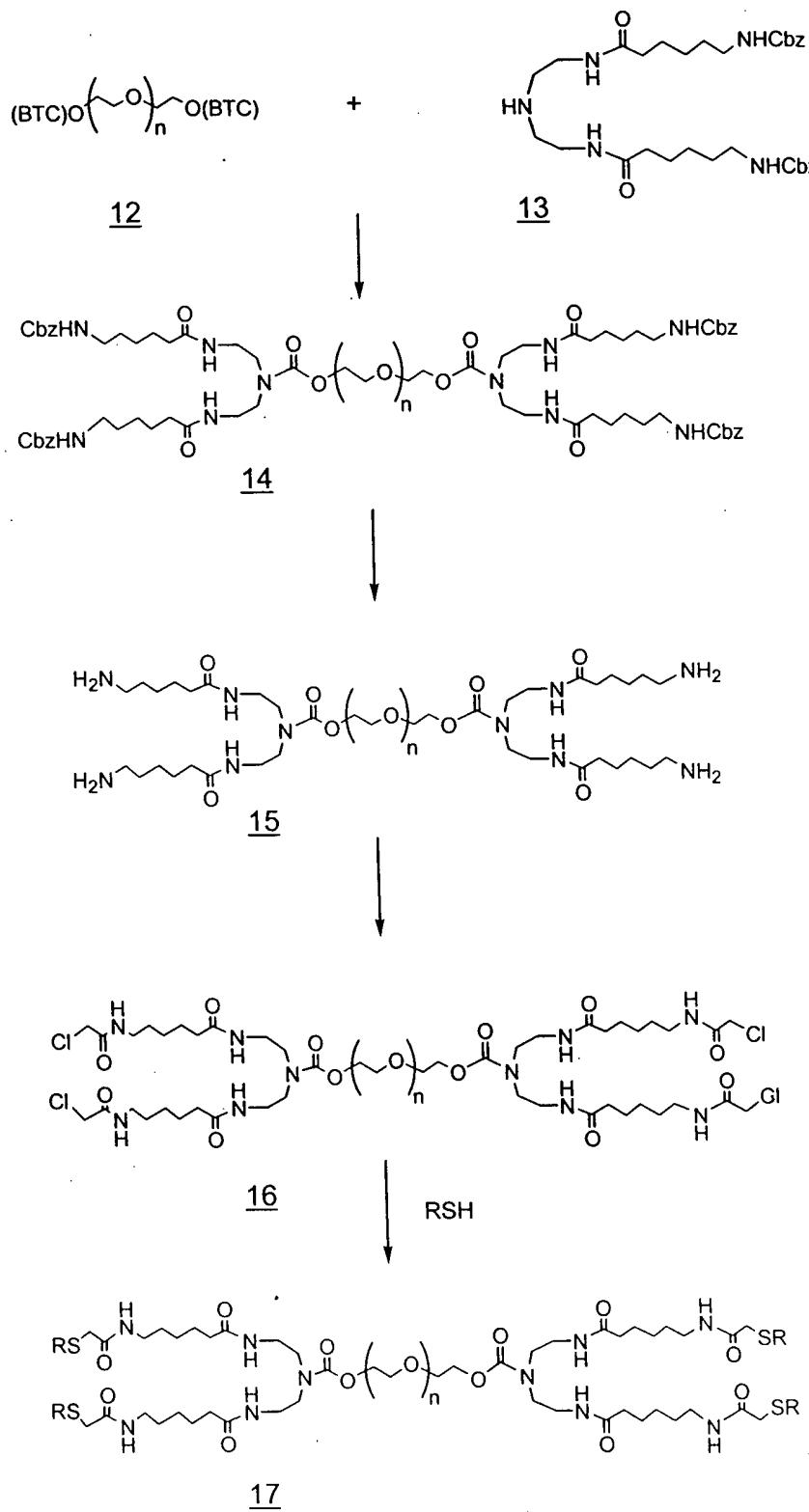


Figure 2

D.G. FIG.		SUBCLASS
CLASS	BY	
DRAFTSMAN		



**Figure 3**

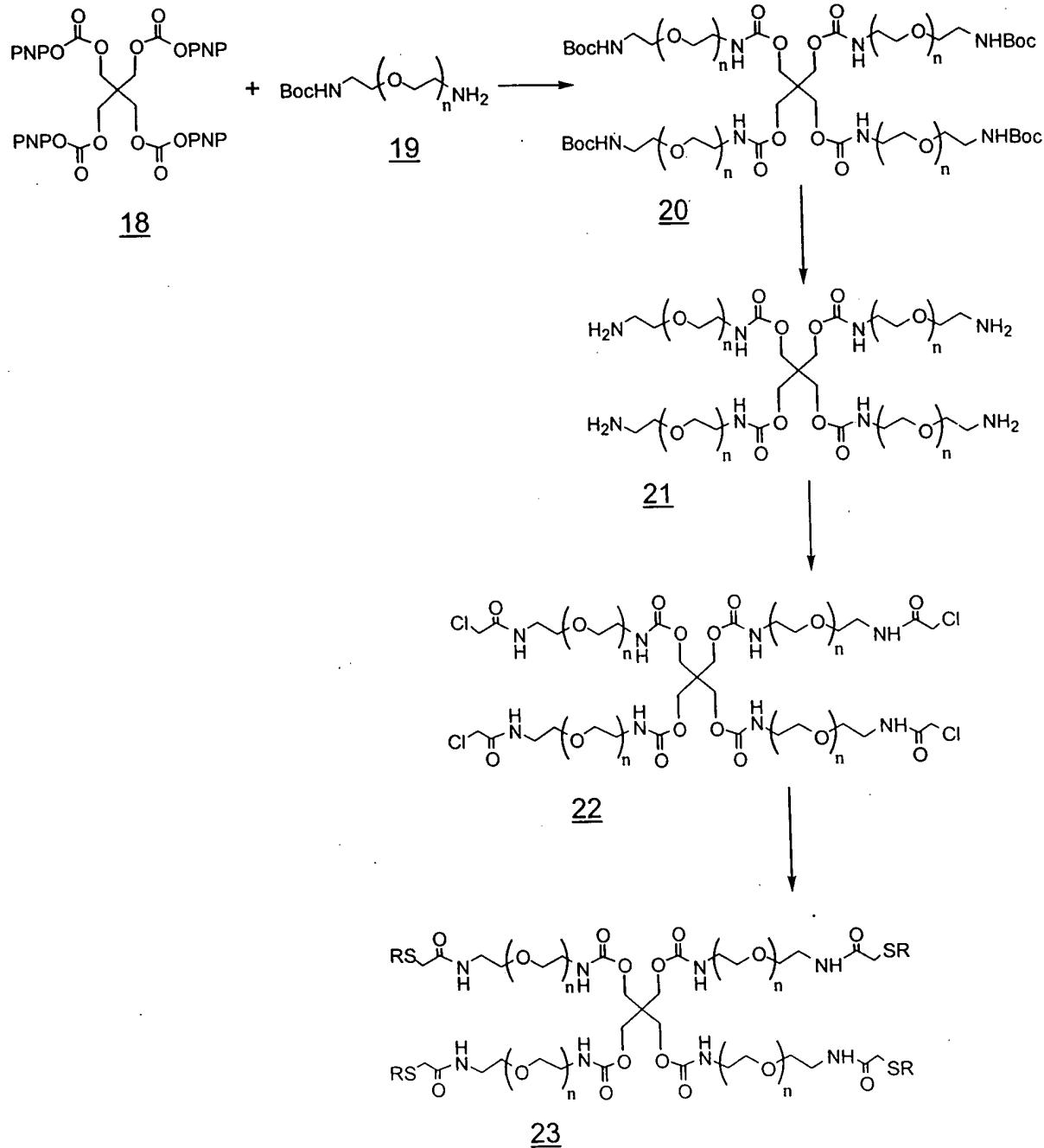


Figure 4

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
CRAFTSMAN		

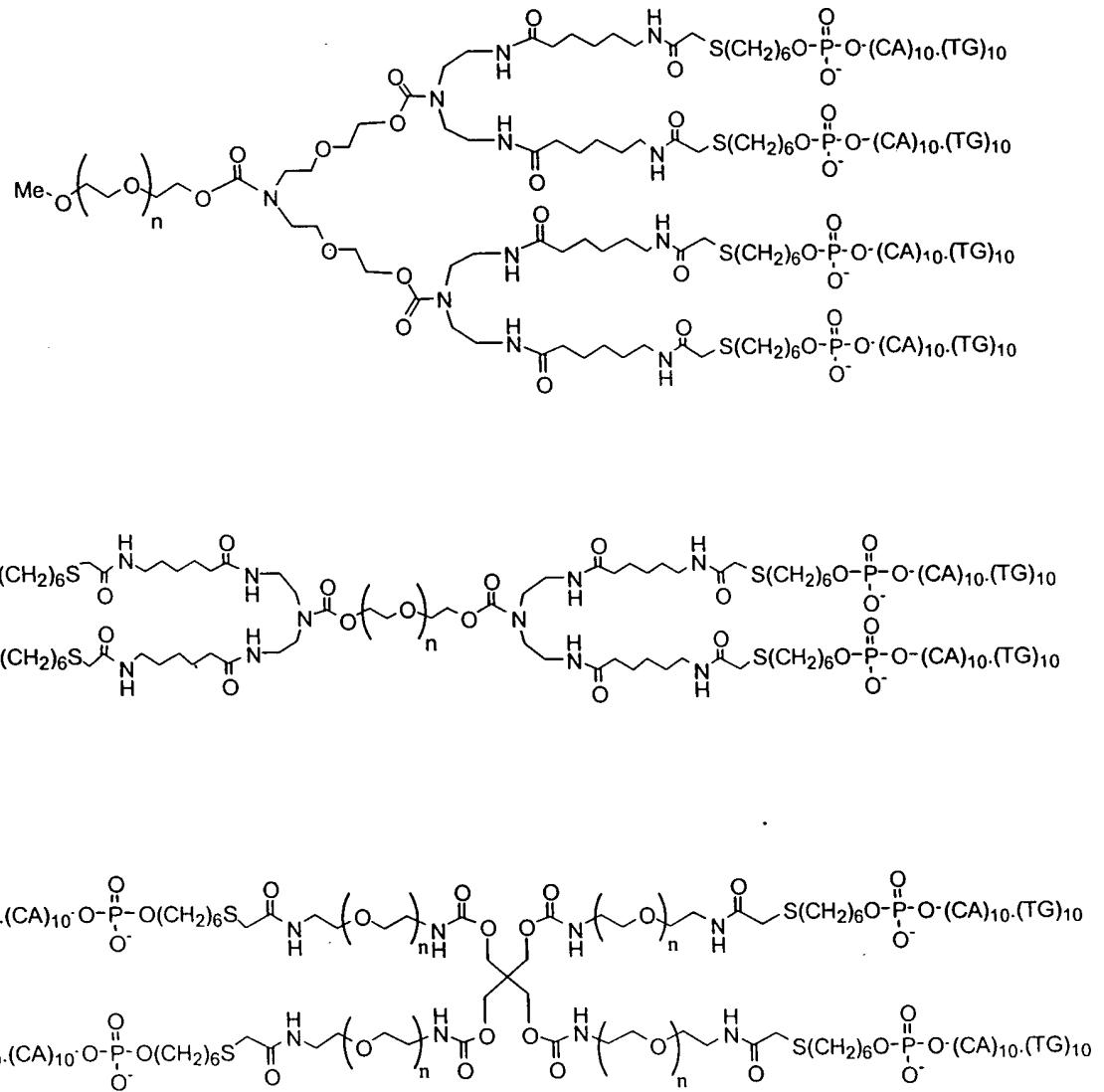


Figure 5

FIG. 6.2. FIG.  
CLASSES SUBCLASS  
GRAFTS:

TECHNIQUE OF PREPARATION

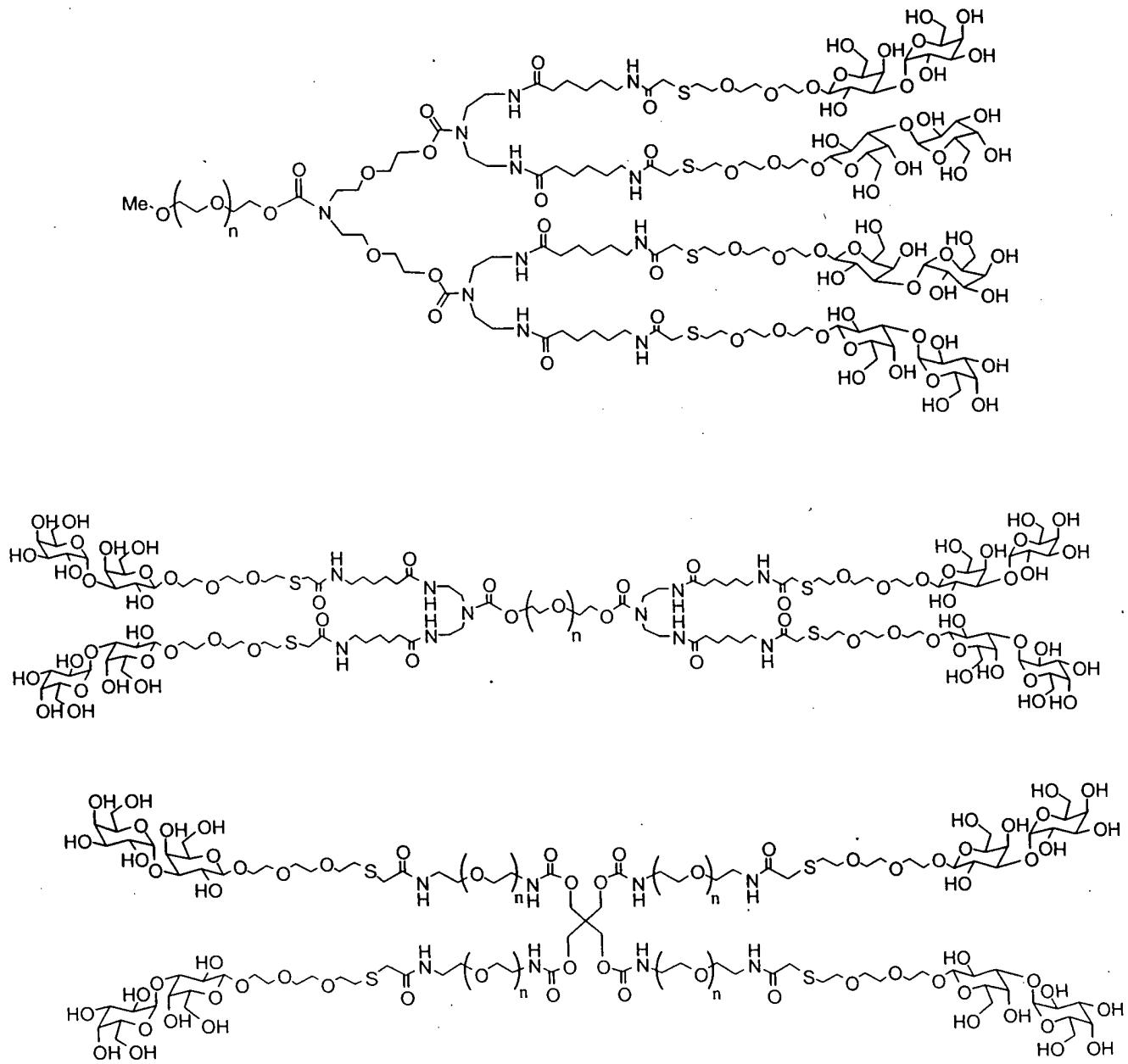
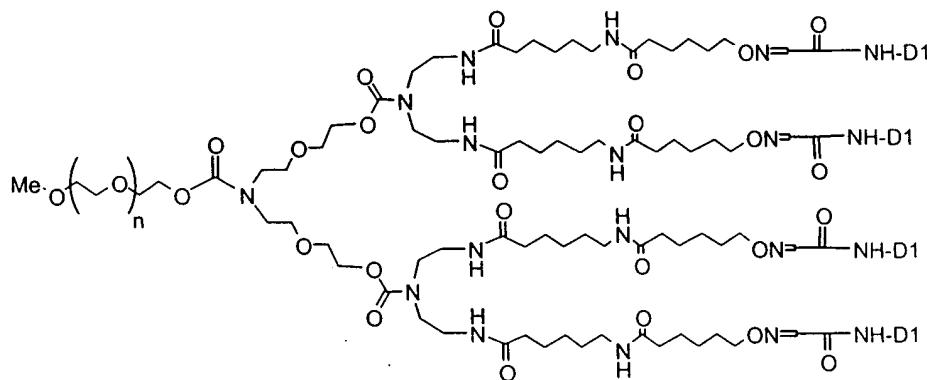
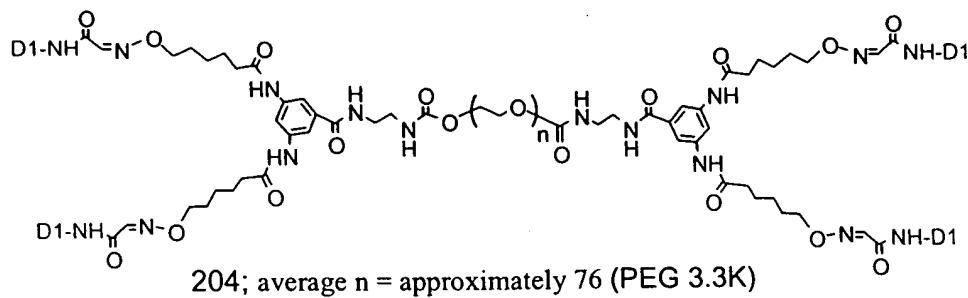


Figure 6

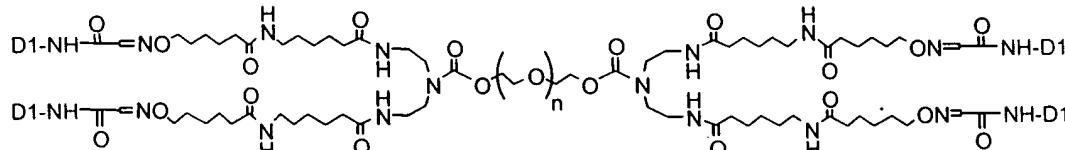


200; average n = approximately 503 (PEG 20K)

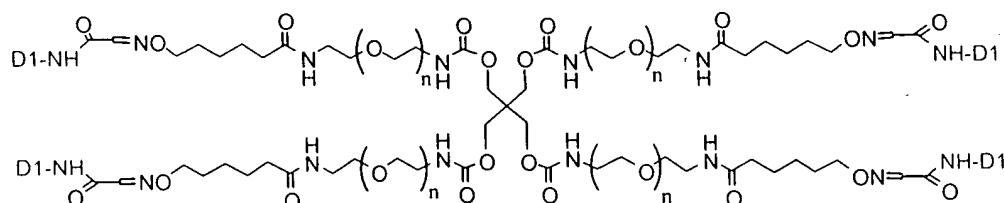
201; average n = approximately 114 (PEG 5K)

205; average n = approximately 261 (PEG 12K)

301; average n = approximately 682 (PEG 30K)



202; average n = approximately 503 (PEG 20K)

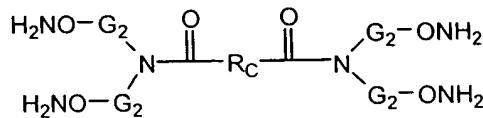


203; average n = approximately 125 (PEG 5K)

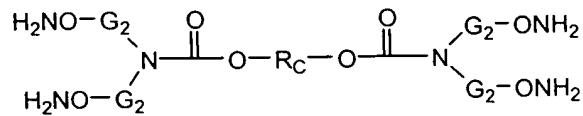
total PEG = 20K

Figure 7

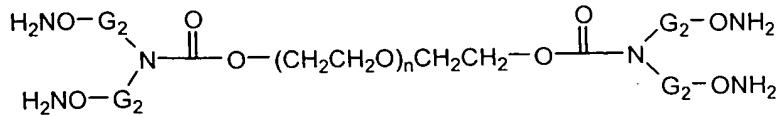
O.C. FIG.	CLASS	SUBCLASS
DP	15	00



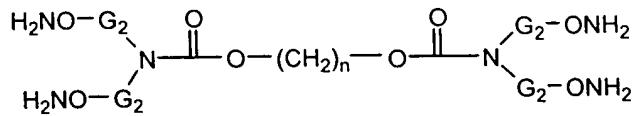
Formula 9



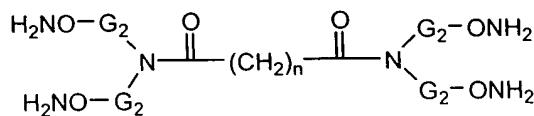
Formula 10



Formula 11



Formula 12



Formula 13

Figure 8

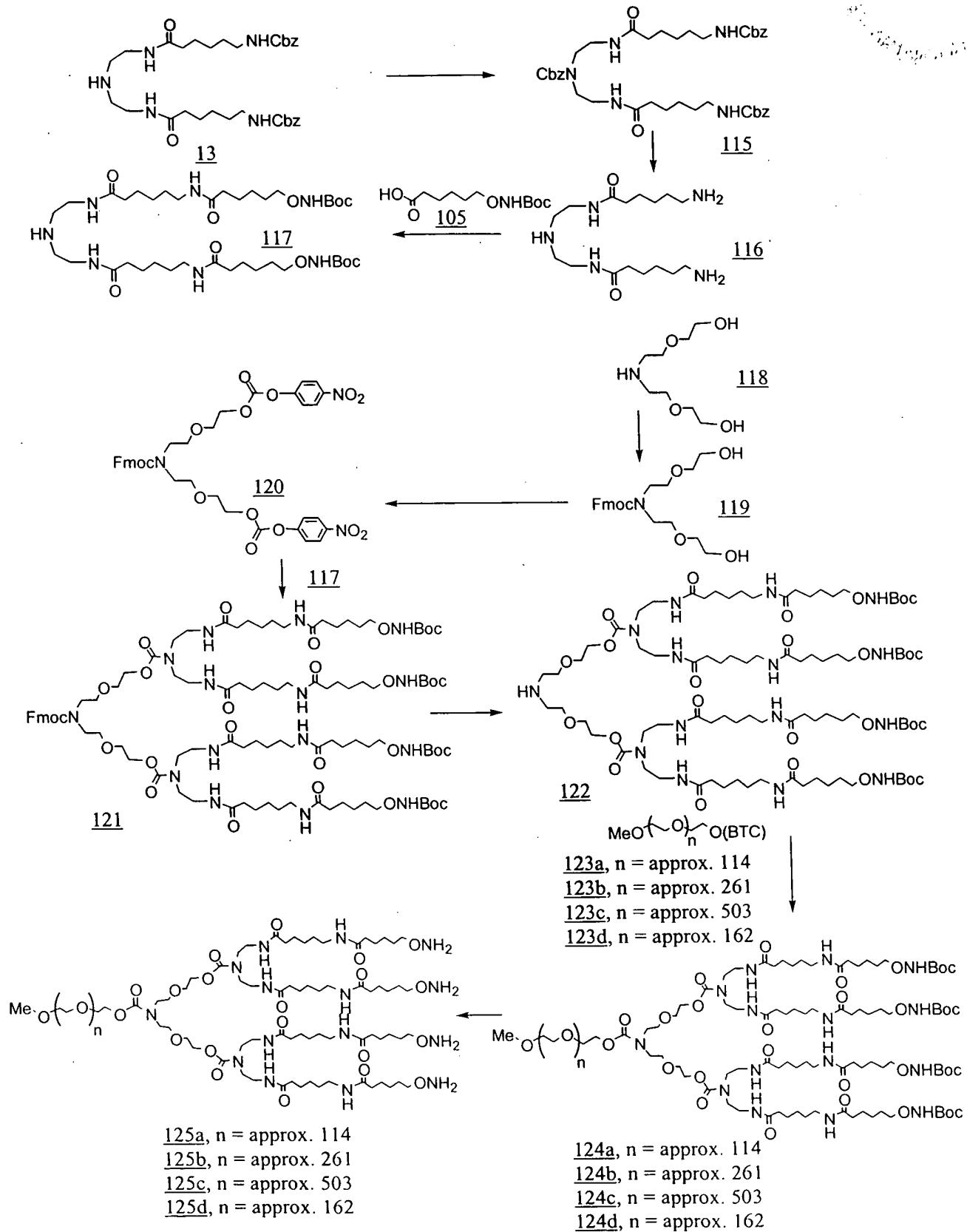


Figure 9

SEARCHED  
 INDEXED  
 COPIED  
 SERIALIZED  
 FILED  
 O.G. FIG.  
 CLASS  
 SUBCLASS  
 DRAFTSMAN

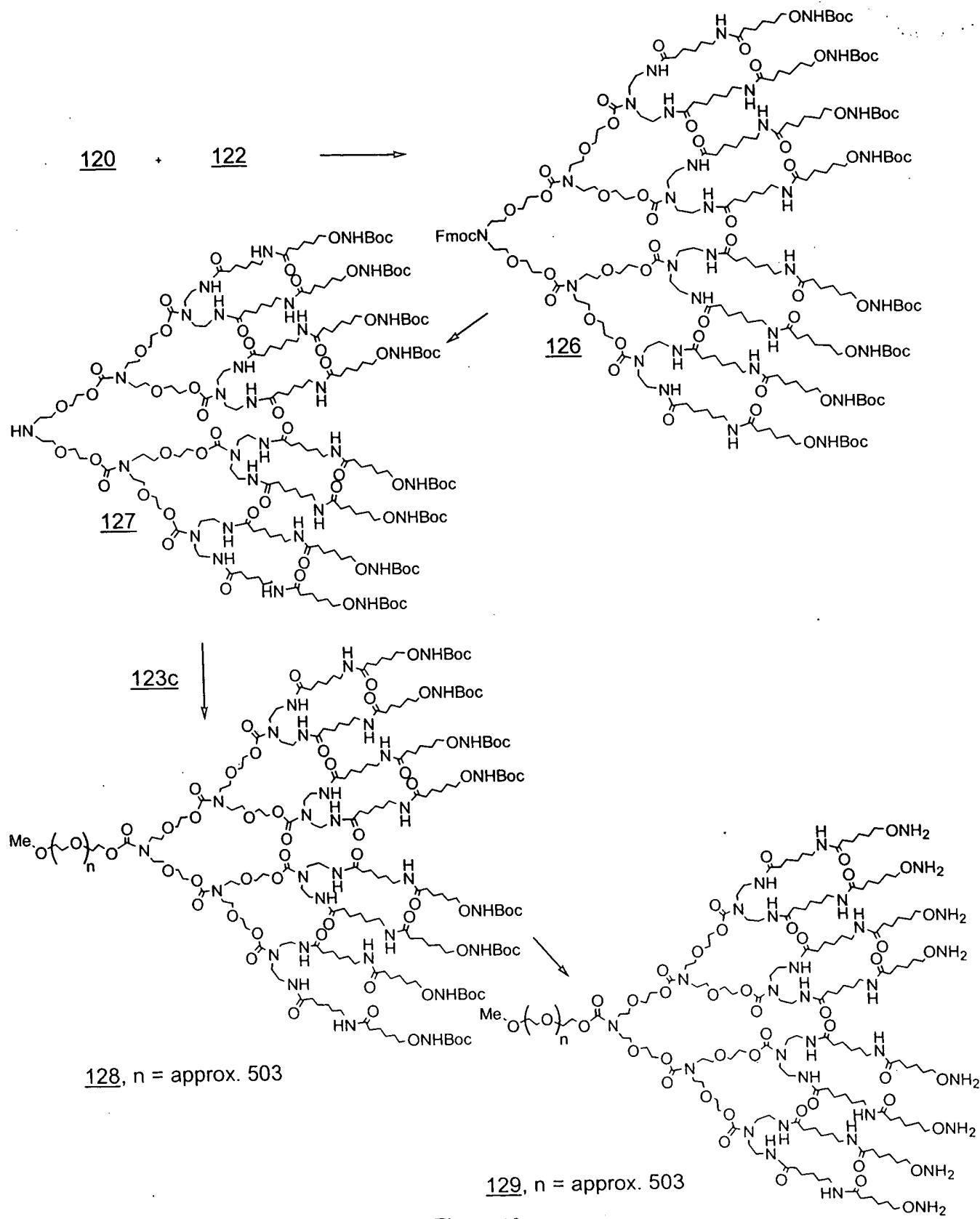


Figure 10

APPROVED BY DRAFTSMAN	O.G. FIG.	CLASS SUBCLASS

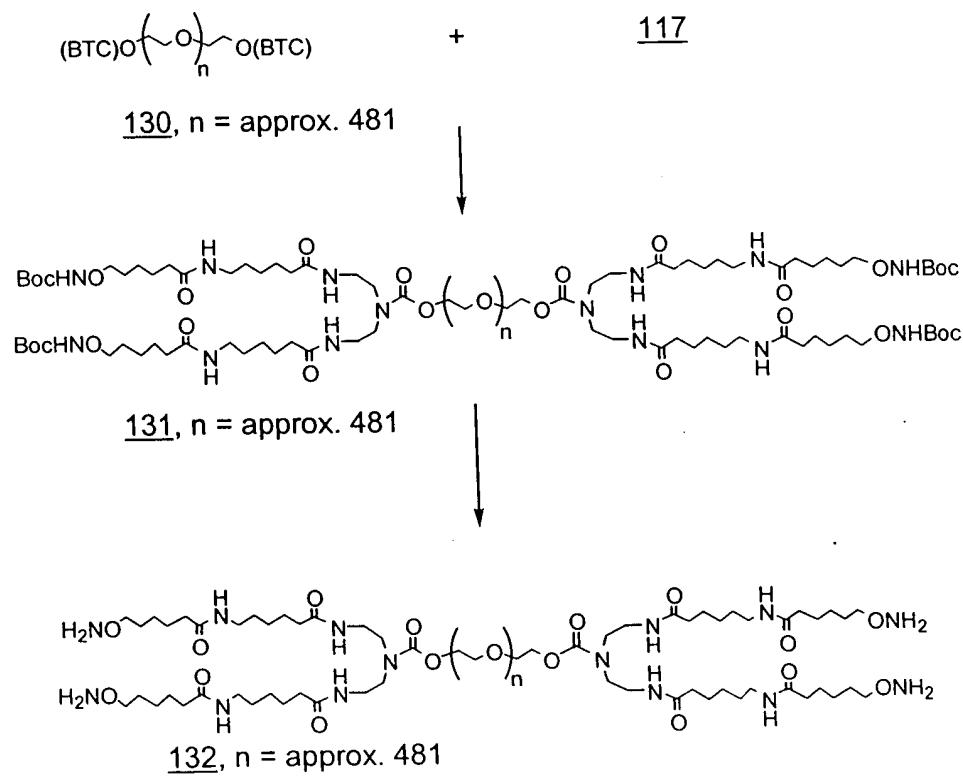


Figure 11

APPROVED BY	O. G. FIG.	CLASS SUBCLASS
	DR. J. W. SMITH	

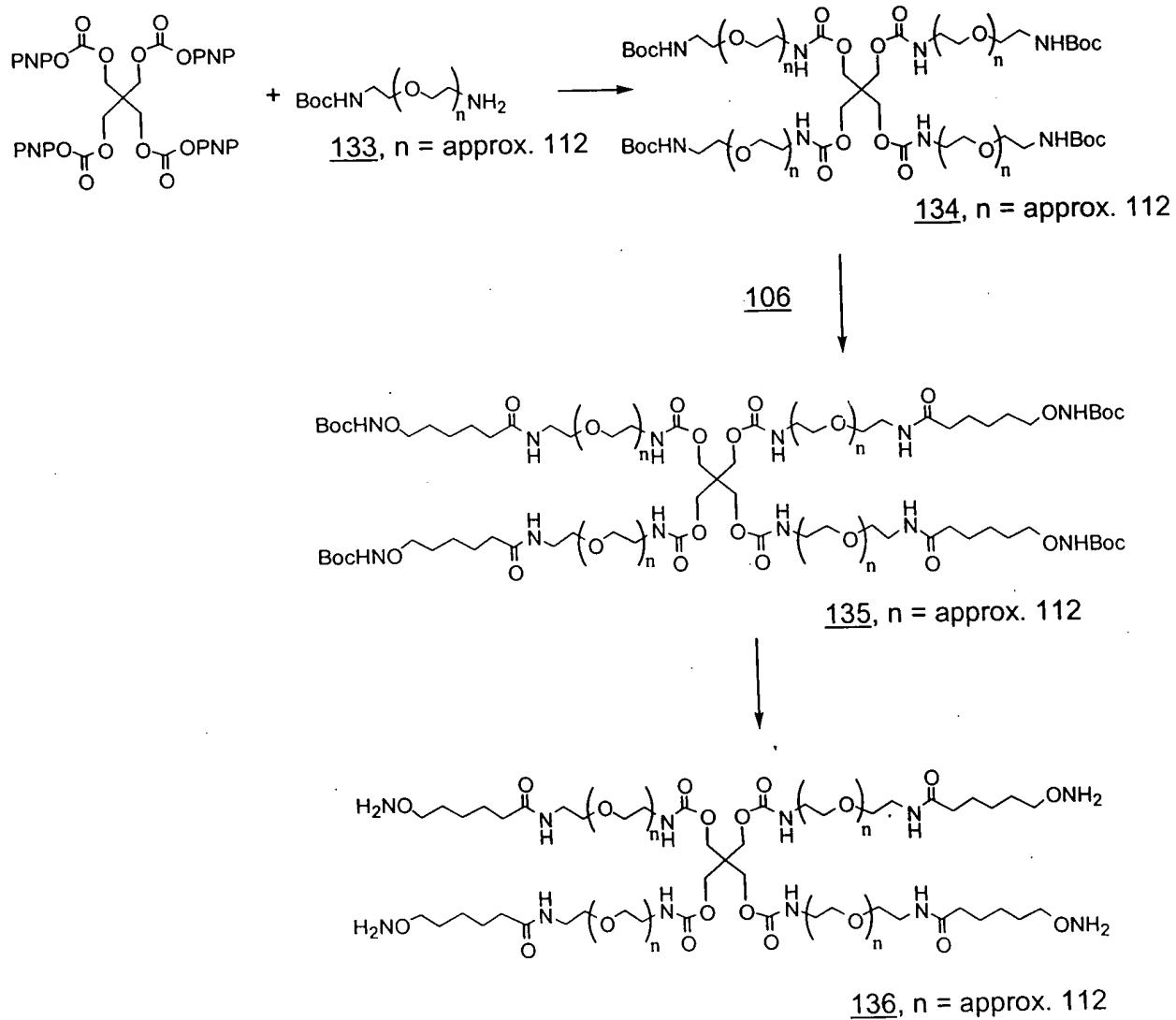


Figure 12

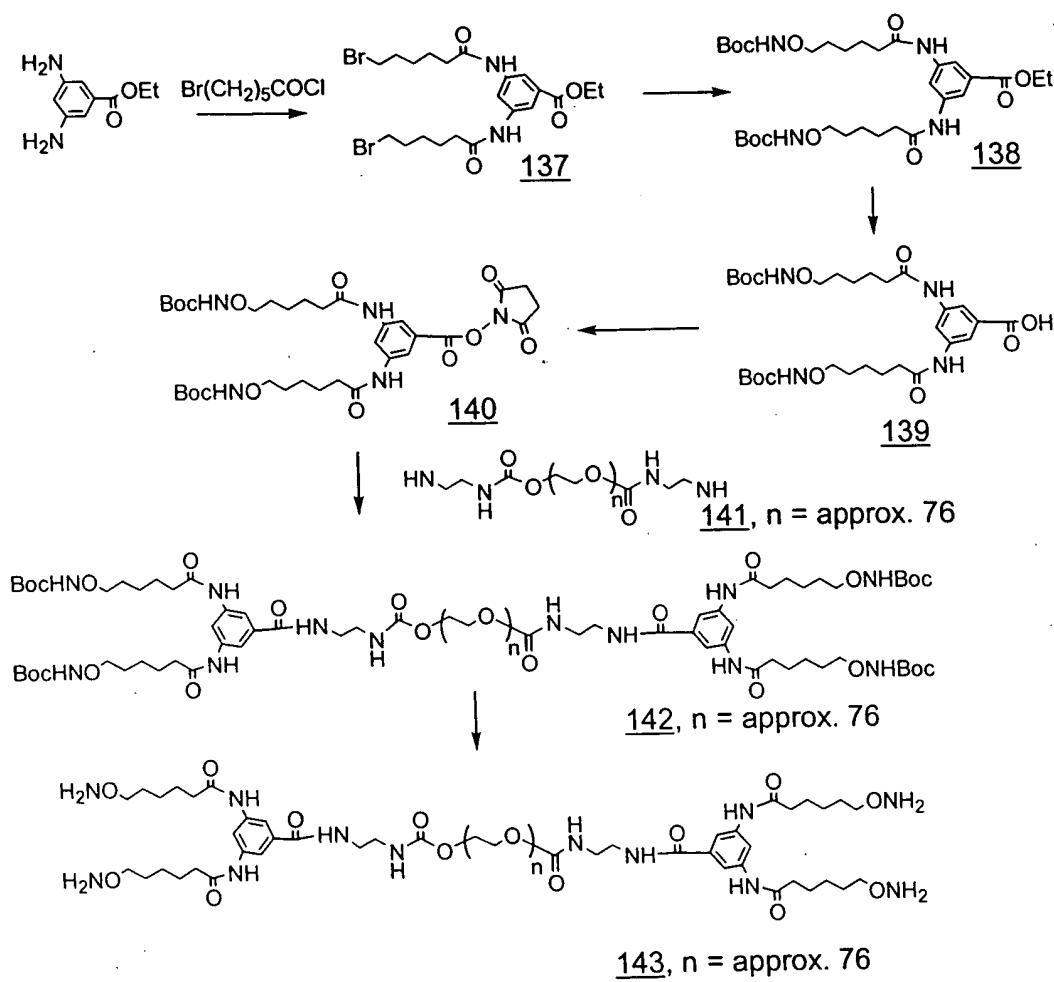


Figure 13

APPROVED	O.C. FIG.	
BY	CLASS	SUBCLASS
D.L. TSIAH		

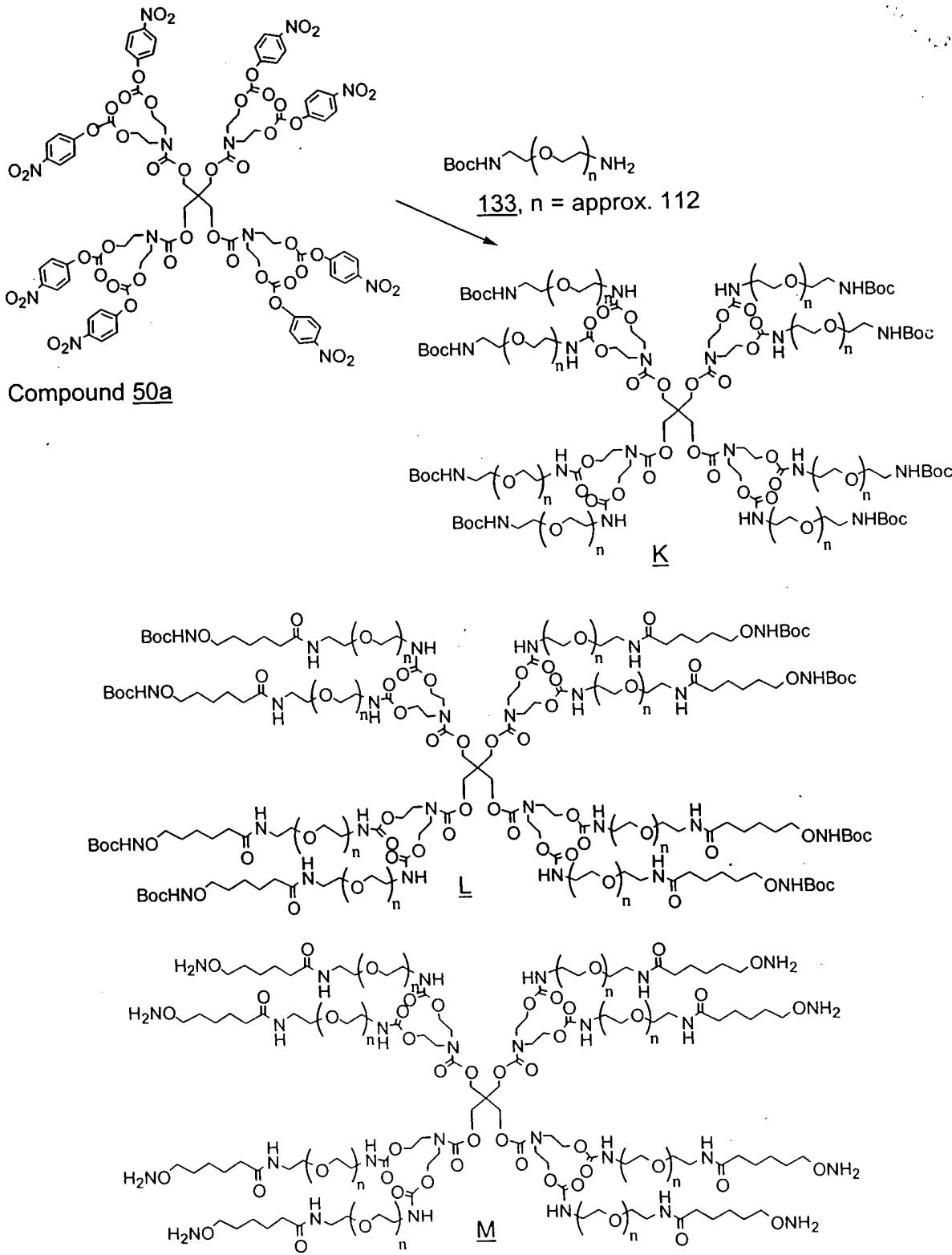


Figure 14

APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

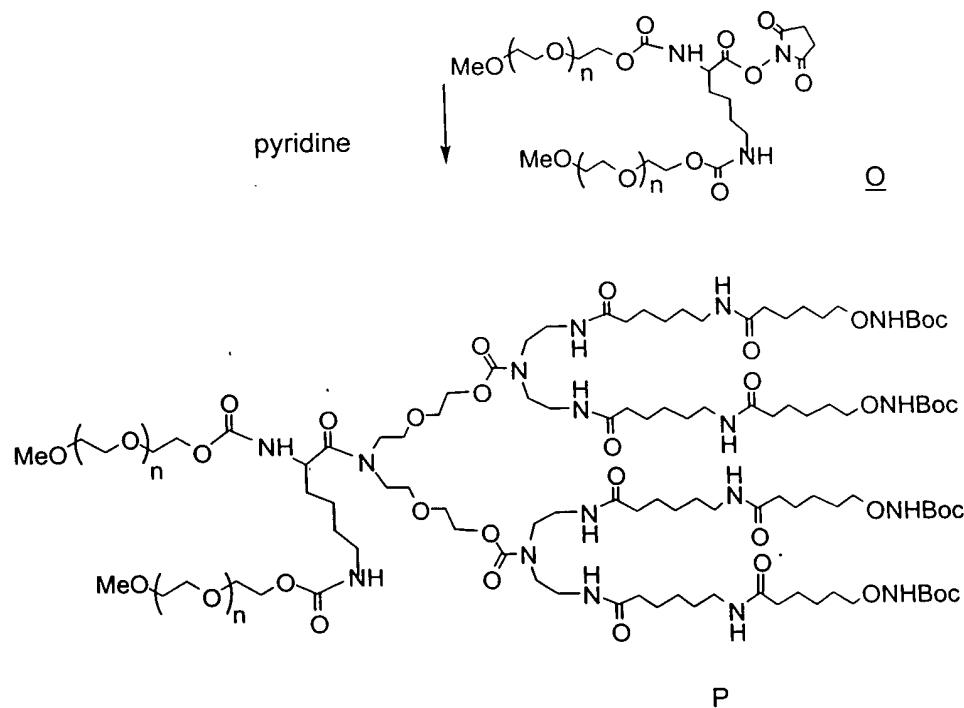
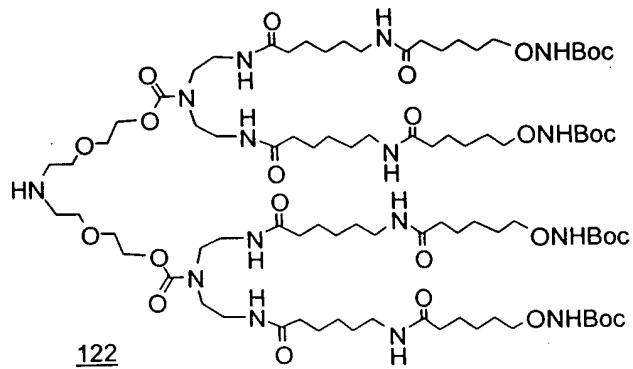


Figure 15

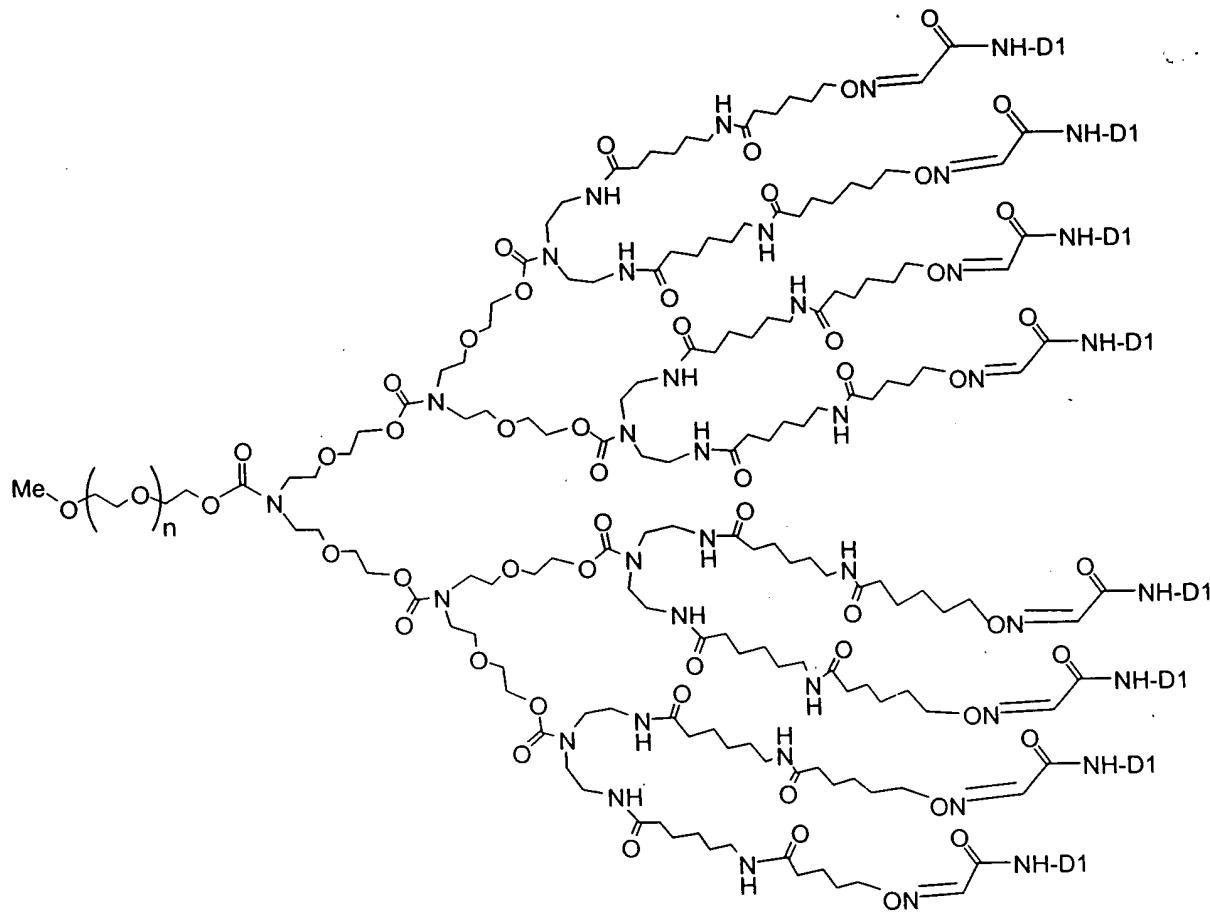


Figure 16

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

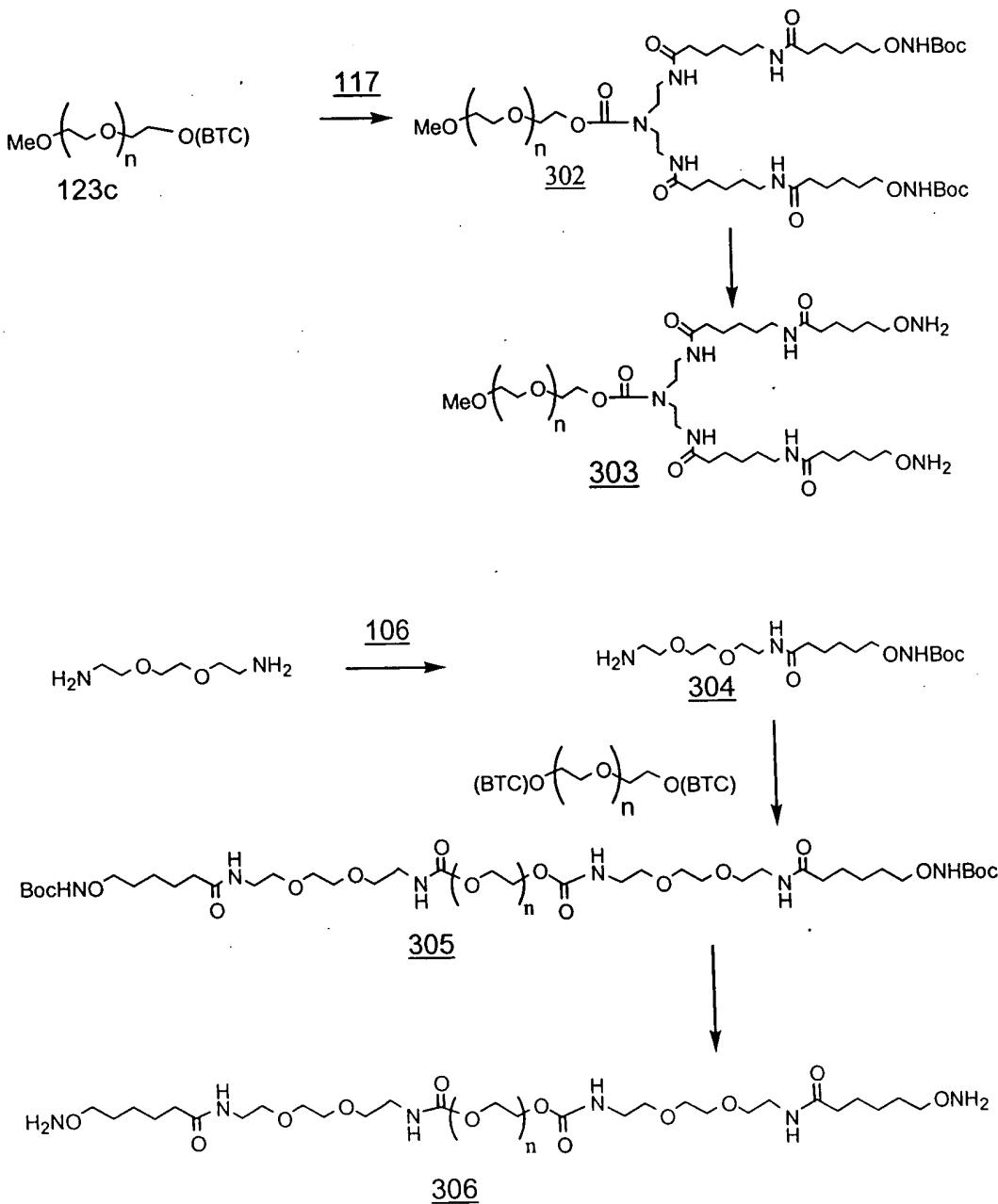


Figure 17

APPROVED	O.G. FIG.	CLASS	SUBCLASS
BY DRAFTSMAN			

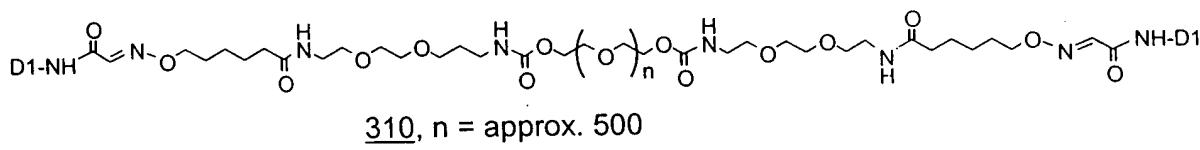
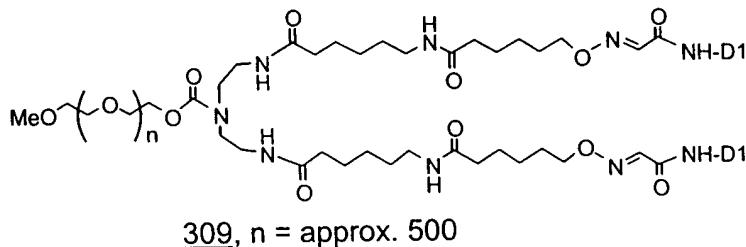


Figure 18

APPROVED	C.G. FIG.
BY	CLASS
DRI-TTSMAN	

gga cgg acc tgt ccc aag cca gat gat tta cca ttt tcc aca gtg gtc 48  
Gly Arg Thr Cys Pro Lys Pro Asp Asp Leu Pro Phe Ser Thr Val Val  
1 5 10 15

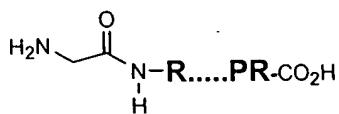
ccg tta aaa aca ttc tat gag cca gga gaa gag att acg tat tcc tgc 96  
Pro Leu Lys Thr Phe Tyr Glu Pro Gly Glu Ile Thr Tyr Ser Cys  
20 25 30

aag ccg ggc tat gtg tcc cga gga ggg atg aga aag ttt atc tgc cct 144  
Lys Pro Gly Tyr Val Ser Arg Gly Gly Met Arg Lys Phe Ile Cys Pro  
35 40 45

ctc aca gga ctg tgg ccc atc aac act ctg aaa tgt aca ccc aga gta 192  
Leu Thr Gly Leu Trp Pro Ile Asn Thr Leu Lys Cys Thr Pro Arg Val  
50 55 60

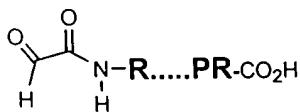
Figure 19

APPROVED	O.G. FIG.	CLASS	SUBCLASS
DRY FTSMAN			



Domain 1 of  $\beta_2$ GPI (D<sub>1</sub>, where bold letters stand for single letter amino acid code of terminal amino acids of Domain 1 of  $\beta_2$ GPI)

pH 5.5  
1-2 M NaOAc  
CuSO<sub>4</sub>  
glyoxylic acid



Transaminated Domain 1 (TAD1)  
Comprising a terminal glyoxyl group

Figure 20